	CRF Processing Date: ///3/20 Changed a file from non-ASCII to ASCIIENTERED CRF Processing Date: ////3/20 Consider the first processing Date: /////3/20 Consider the first processing Date: ////////////////////////////////////
	Changed the margins in cases where the sequence text was "wrapped" down to the next line.
[Edited a format error in the Current Application Data section, specifically:
	Edited the Current Application Data section with the actual current number. The number inputted by the applicant was the prior application data; or other
ř	Added the mandatory heading and subheadings for "Current Application Data".
E	Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
(Changed the spelling of a mandatory field (the headings or subheadings), specifically:
(Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:
1	nserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:
(Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
	Inserted colons after headings/subheadings. Headings edited included:
	Deleted extra, invalid, headings used by an applicant, specifically:
	Deleted: non-ASCII "garbage" at the beginning/end of files; secretary initials/filename at end of file page numbers throughout text; other invalid text, such as
	Inserted mandatory headings, specifically:
	Corrected an obvious error in the response, specifically:
	Edited identifiers where upper case is used but lower case is required, or vice versa.
	Corrected an error in the Number of Sequences field, specifically:
	A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
	Deleted <i>endIng</i> stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error ue to a PatentIn bug). Sequences corrected:
	Other:

*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.



PCT10

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Family
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     To the Sty Sta Ara Va. Asposity Ash As a Sta the Valosty Tyr
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75 - 216 - 282 II 184: 4

-1.5.1F: 11 %:-

11MB: 17:47:4 FAIRNT ARRITARIT N: US/10/069,605 list if if : A:\PTO.AMC.txt .:: .: N:\CRF4\11132002\J069605.raw ". - ..11 - IFHAH: 14 . i. iyik: iki MARKET ROOM BOWER AFTER THE DESIGNATION OF PROPERTY + ... + FLATTEF: ± 1 . The stable interpretability symmetric Equilibries $\pm 1 + 4 + 4 + 19$, while ± 4 55 Wal like Lys War Asy Ite: Wal Ash 310 300 Ata Thr Aly 3th 86 1 89 KAITU CHQ II M: H 90 (211) LENGTH: 14 91 KALI TYFF: 18T 92 /2035 CRGANICM: Artiflet L Degiene 94 KZZZO FRATUFF: 95 (213) (THEF INFOFMATION: Cynthetic Leptide 97 <4565 SEQUENTE: 1 99 Pro Glu The Win Asy The The Tyr Leu Trp Trp Ile Ash Ash. 113 KDIN - 282 II N : x 104 <201 - LENGTH: 09 105 <2125 TYPE: FRT 106 <2135 OPGANISM: Artificial Sequence 108 <226> FEATURE: 109 <203> OTHER INFOFMATION: Synthetic Peptide 101 - 407 / SEQUENCE: A 113 Leu Gin Leu Ser Ash Gly Ash Arg Thr Leu Thr Leu Leu Ser 114 1 116 <2108 SEQ 10 NO: T 117 +311> LENGTH: 14 118 <212> TYPE: FF0 119 <213> ORGANISM: Artificial Sequence 101 (220) PERTURE: 122 <223> OTHER INFORMATION: Synthetic Peptide 124 <400> SEQUENCE: 7 126 Ser Asp Erc Val Thr Leu Ash Val Thr Tyr Gly Fre Asp Thr 127 1 130 <216> SEQ ID NO: 8 131 <211> LENGTH: 14 132 <212> TYFE: PFT 133 <213 > ORGANISM: Artificial Sequence 135 WALL FEATURE: 136 (1663) OTHER INTERNATION: Evenyboria Depaids 138 <400> SEQUENCE: 8 140 Pro Ser Asp Thr Tyr Tyr Ara Pro Gly Ala Ash Lou Ser Leu 141 144 × 216 / SEQ 11 10 : 9 145 - W11 - LENGTH: 14 146 KÜBZ KIYPE: FF1 140 x153 companion: Artificial Sequence 149 · . . t · FEATURE:

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Topical to THEE INFORMATIONS Symiletic English
15° k⊈ ok 15°, MEN N : K
194 Ala Ala Der Alb Bro Bro Ala Shi Iyr Der Irp Deu Leo Ad
18 + +, 1 + 18, 11 N ± 1
18 + +, 11 + 18 N XIB: 14
16 . 1. · 1978: 187
161 - 1152 "FGAMIOM: Artificial Cequence
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164 < 2.3> STHEE INFORMATION: Synthetia Pentiae
166 - 411 > JEQUENTE:
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173 < 1115 LENSTE: 14
 NG KILIK IMPR: BRT
175 Kl13 × CF SAMIRM: Artificial Sequence
179 K21(> FEATURE:
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11MF: 1:4:4:4 : 4 FM1 AFF11 AFF11 N: US/10/069,605 in A:\PTO.AMC.txt :: : N:\CRF4\11132002\J069605.raw * 4 *.. FFAITH: r Electric CHER CAR RMATT Mr Cynthestra Egittir * *4 * **, TFN F: . to the layer only one of the Carl Asp. And Ast. Ast. Ast. Lot 100 (100 (10)) The •1 1 *1* *.1 * JE, 11 W : .1 :14 *.11 * LEMMIE: 14 :15 *.17 * TYPE: :FF 316 külta ku BGANIUM: Artiil mial Sequende 318 - 22 * FEATURE: 319 */Fre THEE INFORMATION: Synthet Do Feptime ALI NA PRANCES AI 323 Val lie lys let Ash Let Met Ser Sid Gld Val ihr Sly Sid 0.1 - FFQ 11 NG: 11 328 4211 - LENGTH: 14 3/9 NOIN - TYPE: FET HBO KUIY WEGANIOM: ArtifildGal Sequence adi. v. 1 × FRAIT Fr: 333 WLLEY CTEFF INFOFMATION: Synthetic Feptide 335 -40 S SEQUENCE: 12 337 Fig. 310 Thr. Win Ash Thr Thr Tyr Leu Trp Trp Val Ash Bly 38 1 341 ×017 × JEG II N : 13 342 (711) LENGTH: 14 343 - XIV - TYFF: FFT 344 - 271 - GEGANIOM: Artificial Sequence 346 FE. G. FEATURE: 347 (223) OTHER INFOFMATION: Synthetic Peptide 349 <400 > SEQUENCE: 13 951 Fie Cer Asy Tim Tyr Tyr Nis Ala Cly Val Ach Leu Ach Leu 352 1 355 <210> SEQ ID NO: 24 356 <011> LENGTH: 14 357 <212> TYPE: FAT 358 KAISH ORGANISM: Artificial Sequence 3€0 <220> FEATUPE: 361 <223> OTHER INFORMATION: Synthetic Peptide 363 ×400> SEQUENTH: 24 R65 Ala Ala Ser Arm Dro Dr. Cor Clm Tyr Ser Trp Dri Mal Ash ((369 <210> SEQ IN NO: 25 376 - 2115 LENGTH: 14 <71 <1100 TYPF: FPT 372 (.13 (CEGANISM: Artificial Sequence :T4 HIZZ FEATURE: *TE - LL: - CTHFF INFO-MATICN: Symiler L: Feyride 377 - 456 - SEQUENCE: 15

47% Ser Val Ash Miv Inr Ele Min Gir Tyr Thr Min Tyr Leo The

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